## Karin Engström

List of Publications by Year in descending order

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279798 434195 1,150 32 23 31 citations h-index g-index papers 32 32 32 1641 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Perfluoroalkyl substances influence DNA methylation in school-age children highly exposed through drinking water contaminated from firefighting foam: a cohort study in Ronneby, Sweden. Environmental Epigenetics, 2022, 8, dvac004.	1.8	11
2	High in Utero Exposure to Perfluoroalkyl Substances from Drinking Water and Birth Weight: A Cohort Study among Infants in Ronneby, Sweden. International Journal of Environmental Research and Public Health, 2022, 19, 2385.	2.6	3
3	Maternal exposure to cadmium during pregnancy is associated with changes in DNA methylation that are persistent at 9Âyears of age. Environment International, 2022, 163, 107188.	10.0	7
4	Serum perfluoroalkyl substances in residents following long-term drinking water contamination from firefighting foam in Ronneby, Sweden. Environment International, 2021, 147, 106333.	10.0	42
5	Early Pregnancy Exposure to Ambient Air Pollution among Late-Onset Preeclamptic Cases Is Associated with Placental DNA Hypomethylation of Specific Genes and Slower Placental Maturation. Toxics, 2021, 9, 338.	3.7	6
6	Associations between serum concentrations of perfluoroalkyl substances and DNA methylation in women exposed through drinking water: A pilot study in Ronneby, Sweden. Environment International, 2020, 145, 106148.	10.0	21
7	Association between serum concentrations of perfluoroalkyl substances (PFAS) and expression of serum microRNAs in a cohort highly exposed to PFAS from drinking water. Environment International, 2020, 136, 105446.	10.0	44
8	Maternal Long-Chain Polyunsaturated Fatty Acid Status, Methylmercury Exposure, and Birth Outcomes in a High-Fish-Eating Mother–Child Cohort. Journal of Nutrition, 2020, 150, 1749-1756.	2.9	5
9	Polymorphisms in potential mercury transporter ABCC2 and neurotoxic symptoms in populations exposed to mercury vapor from goldmining. Environmental Research, 2019, 176, 108512.	7.5	15
10	Association of Arsenic Exposure with Whole Blood DNA Methylation: An Epigenome-Wide Study of Bangladeshi Adults. Environmental Health Perspectives, 2019, 127, 57011.	6.0	40
11	Maternal polymorphisms in glutathione-related genes are associated with maternal mercury concentrations and early child neurodevelopment in a population with a fish-rich diet. Environment International, 2018, 115, 142-149.	10.0	34
12	Associations between Methylated Metabolites of Arsenic and Selenium in Urine of Pregnant Bangladeshi Women and Interactions between the Main Genes Involved. Environmental Health Perspectives, 2018, 126, 027001.	6.0	10
13	Prenatal arsenic exposure is associated with increased plasma IGFBP3 concentrations in 9-year-old children partly via changes in DNA methylation. Archives of Toxicology, 2018, 92, 2487-2500.	4.2	33
14	Arsenic exposure from drinking water is associated with decreased gene expression and increased DNA methylation in peripheral blood. Toxicology and Applied Pharmacology, 2017, 321, 57-66.	2.8	37
15	CYP3A genes and the association between prenatal methylmercury exposure and neurodevelopment. Environment International, 2017, 105, 34-42.	10.0	24
16	Transcriptomics and methylomics of CD4-positive T cells in arsenic-exposed women. Archives of Toxicology, 2017, 91, 2067-2078.	4.2	26
17	AS3MT-mediated tolerance to arsenic evolved by multiple independent horizontal gene transfers from bacteria to eukaryotes. PLoS ONE, 2017, 12, e0175422.	2.5	29
18	Exposure to Inorganic Arsenic Is Associated with Increased Mitochondrial DNA Copy Number and Longer Telomere Length in Peripheral Blood. Frontiers in Cell and Developmental Biology, 2016, 4, 87.	3.7	42

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19	Polymorphisms in ATP-binding cassette transporters associated with maternal methylmercury disposition and infant neurodevelopment in mother-infant pairs in the Seychelles Child Development Study. Environment International, 2016, 94, 224-229.	10.0	32
20	Genetic variation in FADS genes is associated with maternal long-chain PUFA status but not with cognitive development of infants in a high fish-eating observational study. Prostaglandins Leukotrienes and Essential Fatty Acids, 2015, 102-103, 13-20.	2.2	34
21	Gene-Environment Interactions for Metals. , 2015, , 239-264.		3
22	Prenatal lead exposure is associated with decreased cord blood DNA methylation of the glycoprotein VI gene involved in platelet activation and thrombus formation. Environmental Epigenetics, 2015, 1, dvv007.	1.8	28
23	Selenium metabolism to the trimethylselenonium ion (TMSe) varies markedly because of polymorphisms in the indolethylamine N-methyltransferase gene. American Journal of Clinical Nutrition, 2015, 102, 1406-1415.	4.7	40
24	The effects of arsenic exposure on blood pressure and early risk markers of cardiovascular disease: Evidence for population differences. Environmental Research, 2015, 140, 32-36.	7.5	31
25	Human Adaptation to Arsenic-Rich Environments. Molecular Biology and Evolution, 2015, 32, 1544-1555.	8.9	113
26	Polymorphisms in ABC Transporter Genes and Concentrations of Mercury in Newborns – Evidence from Two Mediterranean Birth Cohorts. PLoS ONE, 2014, 9, e97172.	2.5	39
27	Polymorphisms in Genes Encoding Potential Mercury Transporters and Urine Mercury Concentrations in Populations Exposed to Mercury Vapor from Gold Mining. Environmental Health Perspectives, 2013, 121, 85-91.	6.0	54
28	N-6-Adenine-Specific DNA Methyltransferase $1$ ( $\langle i \rangle$ N6AMT1 $\langle i \rangle$ ) Polymorphisms and Arsenic Methylation in Andean Women. Environmental Health Perspectives, 2013, 121, 797-803.	6.0	40
29	Possible Positive Selection for an Arsenic-Protective Haplotype in Humans. Environmental Health Perspectives, 2013, 121, 53-58.	6.0	44
30	Pregnancy and the methyltransferase genotype independently influence the arsenic methylation phenotype. Pharmacogenetics and Genomics, 2012, 22, 508-516.	1.5	28
31	Arsenic Exposure through Drinking Water Is Associated with Longer Telomeres in Peripheral Blood. Chemical Research in Toxicology, 2012, 25, 2333-2339.	3.3	79
32	Polymorphisms in Arsenic(+III Oxidation State) Methyltransferase ( <i>AS3MT</i> ) Predict Gene Expression of <i>AS3MT</i> as Well as Arsenic Metabolism. Environmental Health Perspectives, 2011, 119, 182-188.	6.0	156